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730 AAAACAGCGGCGAGA 744
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SEQUENCE ?
US-09-431-404-1525/c
Sequence 1525, Application US/09491404
GENERAL NAME: Yunnan T.
APPLICANT: Tian Xunhua T.
APPELLANT: Tillinghaert, John
APPELLEE: Sianku, Ankura
APPELLOUT: Liao Shengyue
APPLICANT: Duanmanc, Radjoje T.
TITLE OF INVENTION: Novel Contigs Obtained From Various Larvae
TITLE OF INVENTOR: From various Larvae
CURRENT APPLICATION NUMBER: US/09/491,404
CURRENT FILING DATE: 2000-01-27
OTHER INFORMATION:
SOFTWARE: pr SP_genes Version 1.0
SEQ ID NO 1525
LENGTH: 866
ORGANISM: Homo sapiens
FEATURES:
LOCATION: 166 bp peptide
LOCATION: (70) ... (205)
OTHER INFORMATION: This location contains the signal peptide sequence,
OTHER INFORMATION: KMTNMLVTHALLQALVGVTSERENSPCHALLSEPTLEQGS, Run with SignalP
NAME/KEY: misc feature
LOCATION: (34)...(700)
OTHER INFORMATION: Similar to G51Y0139 in the geneprep database release 114,
OTHER INFORMATION: Run with PASTYT 3.3000, default parameters
US-09-431-404-1525
Alignment Scores:
Prod. No.: 0 Length: 886
Score: 55.00 Conservative: 15
Percent Identity: 25.0%
Sequence Similarity: 25.0%
Query Match: 4.6%
DB: 5
Gap#: 1
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US-10-033-958-61-COPY x US-09-491-404-1525 (1-886)
64 CysHemAmrPygAglyvylIhFtsErTpmcUpGloPrIeValIySpHeRodly 83
617 TCCTTCAGGTCCGCCGGTGGAAGC...GTTCGAGAAATCTCTTTGAGA 594
84 AlvalApagAlaThrytArIdaLeMevAlaProPcpAlaProGrAsGa 101
AlvalApagAlaThrytArIdaLeMevAlaProPcpAlaProGrAsGa 101
593 GCC...TCGATTGTTTTTCTCGGAGGAGCGAGCATCTCTTCCT 543
104 GlusPArgAlaGspRhpTpaqHlAtLeValTrApLleTyAlaAspLeu 123
GCATCAATCAAGCAACTACTTAATACCCCTGG...NAGCATCATGTCG 498
124 LyalwylDylAladIlLydInGu...
497 CGTGCGGGAGGAGCGCTCGTAGGCTGTACTCGCTCGTAATCTCTTCCTTC 148
LeiderAlArtydAlaAPASpGspFrphAlHlsSeCyPhpe 317
437 AGSTCGGCGCCTTGAATCT...TACCACCAATCTCTCTGAGTCTCTGGATTC 379
133 .....HlsAsPygdln 151
446 .....
378 TGCTGTCTAGSGCATCTGATCACCATCAAG 343
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Search completed: June 23, 2006, 15:58:13
0.001 gpc#

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Search completed: June 23, 2006, 15:58:13
Job time : 0.001 secs